

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING

ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _____

Source: _____

Date Processed by STIC: _____

10/563,277
IFWP
02/08/2007

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

DATE: 02/08/2007

PATENT APPLICATION: US/10/563,277

TIME: 11:49:45

Input Set : A:\PTO.SS.txt

Output Set: N:\CRF4\02082007\J563277.raw

3 <110> APPLICANT: Institut National de la Recherche Agronomique (INRA)

5 <120> TITLE OF INVENTION: Method of producing double low restorer lines of Brassica
napus

6 having a good agronomic value

8 <130> FILE REFERENCE: D21413

C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/563,277

C--> 10 <141> CURRENT FILING DATE: 2006-01-04

E--> 10 <160> NUMBER OF SEQ ID NOS: 5

12 <170> SOFTWARE: PatentIn version 3.2

found 14 Does Not Comply
Corrected Diskette Needed
(pg -1)

ERRORED SEQUENCES

230 <210> SEQ ID NO: 14

231 <211> LENGTH: 21

232 <212> TYPE: DNA

233 <213> ORGANISM: artificial sequence

235 <220> FEATURE:

236 <223> OTHER INFORMATION: pCP418 L primer

238 <400> SEQUENCE: 14

239 aatttctcca tcacaaggac c

21

E--> 249 4

deleted

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/563,277

DATE: 02/08/2007

TIME: 11:49:46

Input Set : A:\PTO.SS.txt

Output Set: N:\CRF4\02082007\J563277.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:249 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:10 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (5) Counted (14)